

<400> 3

gagagaga

SEQUENCE LISTING <110> Dim <120> METHODS FOR DETECTION AND QUANTIFICATION OF ANALYTES IN COMPLEX MIXTURES <130> 11616-004-999 <140> 09/898,743 <141> 2001-07-03 <160> 3 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 20 <212> DNA <213> Artificial Sequence <220> <221> repeat\_unit <222> (1)...(20)  $\langle 223 \rangle$  repeats n times, n=1,2,3,4,5,6,7,8,9,10 <400> 1 actctctct tctctctc 20 <210> 2 <211> 20 <212> DNA <213> Artificial Sequence <220> <221> repeat unit <222> (1)...(20)  $\langle 223 \rangle$  repeats m times, m=1,2,3,4,5,6,7,8,9,10 gctctctct tctctctc 20 <210> 3 <211> 10 <212> DNA <213> Artificial Sequence <220> <223> primer

10